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The new GeneChip® One-Cycle and Two-Cycle cDNA Synthesis Kits.


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-> all probe sets (7129)
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Full Record

Details for HUGENEFL:L17131_RNA1_AT

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GeneChip Array Information

Probe Set ID L17131_rna1_at
GeneChip Array HumanGeneFL Array
Organism Common Name Human

Probe Design Information

Transcript ID L17131_rna1
Sequence Type Exemplar sequence
Representative Public ID L17131 [NCBI](#)
Target Description L17131, class A, 20 probes, 20 in L17131mRNA#1 1646-2198, Human high mobility group protein (HMG-I(Y)) gene exons 1-8, complete cds

Sequence

>HUGENEFL:L17131_RNA1_AT
ttgtccagggtgaggcccaagagccctgtggccgccacctgaggtgggctggggctgctcc
cctaaccctactttcgttcgccactcagccatttccccctcctcagatggggcaccaat
aacaaggagctcaccctgccgctcccaacccccctcctgctcctcctgcccccaagg
ttctggttccatttttccctctgttcacaaactacctctggacagtgtgtgtgtttttgt
tcaatgttccattcttcgacatccgtcattgtctgctgctaccagcgccaaatgttcaccc
tcattgcctcctgttctgcccacgatccctcccccagatactctttgtggggaagagg
ggctggggcatggcaggctgggtgaccgactacccagtcaccaggaagggtggggccctg
cccctaggatgctgcagcagagtgcagcaagggggcccaatcgaccataaagggtgtagg
ggccacctcctccccctgttctgttggggaggggtagccatgatttgtcccagcctgggg
ctccctctctggtttcctatttgcagttacttgaata

Target Sequence

Probe Sequence(5'-3')	Probe		Probe	Strandedness
	X	Y	Interrogation Position	
TTGTCCAGGTGAGGCCCAAGAGCCC	294	101	1658	Antisense
AGGTGAGGCCCAAGAGCCCTGTGGC	295	101	1664	Antisense
ACCAATAACAAGGAGCTCACCTGC	296	101	1772	Antisense
TTTTCTCTGTTACAAACTACCTC	297	101	1850	Antisense
CTACCTCTGGACAGTTGTGTTGTTT	298	101	1868	Antisense
TTCCATTCTTCGACATCCGTCATTG	299	101	1904	Antisense
TCTTCGACATCCGTCATTGCTGCTG	300	101	1910	Antisense

Probe Info	GCTACCAGCGCCAAATGTTTCATCCT	301	101	1934	Antisense
	TCATCCTCATTGCCTCCTGTTCTGC	302	101	1952	Antisense
	TCATTGCCTCCTGTTCTGCCCACGA	303	101	1958	Antisense
	AAGATACTCTTTGTGGGGAAGAGGG	304	101	1994	Antisense
	GCAGGCTGGGTGACCGACTACCCCA	305	101	2030	Antisense
	CCCCTAGGATGCTGCAGCAGAGTGA	306	101	2078	Antisense
	AGCAAGGGGGCCCGAATCGACCATA	307	101	2102	Antisense
	CGAATCGACCATAAAGGGTGTAGGG	308	101	2114	Antisense
	GCCATGATTTGTCCCAGCCTGGGGC	309	101	2174	Antisense
	CTGGGGCTCCCTCTCTGGTTTCCTA	310	101	2192	Antisense
	CTCCCTCTCTGGTTTCCTATTTGCA	311	101	2198	Antisense
	CTCTGGTTTCCTATTTGCAGTTACT	312	101	2204	Antisense
	TTTCCTATTTGCAGTTACTTGAATA	313	101	2210	Antisense

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